



1600

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/502,945

DATE: 03/06/2003

TIME: 16:14:35

Input Set : N:\Crf3\RULE60\09502945.raw
 Output Set: N:\CRF4\03062003\I502945.raw

17
17

1 <110> APPLICANT: Scanlan, Matthew J.
 2 Chen, Yao-Tseng
 3 Stockert, Elisabeth
 4 Old, Lloyd J.
 5 <120> TITLE OF INVENTION: ISOLATED NUCLEIC ACID MOLECULES
 6 ASSOCIATED WITH COLON CANCER AND METHODS FOR DIAGNOSING AND
 7 TREATING COLON CANCER
 8 <130> FILE REFERENCE: LUD-5506-JEL/NDH
 9 <140> CURRENT APPLICATION NUMBER: 09/502,945
 10 <141> CURRENT FILING DATE: 2000-02-11
 11 <150> PRIOR APPLICATION NUMBER: US/08/948,705
 12 <151> PRIOR FILING DATE: 1997-10-10
 13 <160> NUMBER OF SEQ ID NOS: 8
 14 <170> SOFTWARE: FastSEQ for Windows Version 3.0
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 16 <211> LENGTH: 1552
 17 <212> TYPE: DNA
 18 <213> ORGANISM: Homo Sapiens
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 23 aaagagcaac taaagcataa agaatttctt ctggctgcta atacttgtaa ccgtgttgg 180
 24 ggtctttgtt tgaaatgtgc tcagcatgaa gctgttcttt cccaaaccca tactaatgtt 240
 25 catatgcaga ccatcgaaag actggtaaaa gaaagagatg acttgatgtc tgcactagtt 300
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 30 agggagtaca tggatcaaa gatgttgc ttgtctcaga atattgccc actggaggcc 600
 31 caggtggaaa aggttacaaa gaaaaagatt tcagctatta atcaactgga gaaaattcaa 660
 32 agccagctgg cttctcgaaa aatggatgtc acaaagggtgt gtggagaaat gcgctatcag 720
 33 ctgaataaaa ccaacatgga gaaggatgag gcagaaaagg agcacagaga gttcaagagca 780
 34 aaaactaaca gggatcttga aattaaagat cagggaaatag agaaaatttgaa aatagaactg 840
 35 gatgaaagca aacaacactt ggaacaggag cagcagaagg cagccctggc cagagaggag 900
 36 tgcctgagac taacagaact gctggcgaa tctgagcacc aactgcaccc caccagatct 960
 37 gaaatagctc aactcagtc agaaaaagg tatacatatg ataaattggg aaagtacag 1020
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 40 ctcctcagca agcagaacca gcttctcctg gagaggcaga gcctgtcgga agaggtggac 1200
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 43 caacgcacca cgaccttccc agggtgacac cgcctcagcc tgcagtgggg ctggccctca 1380
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45 gcagactcct ggttatgtt ttcagaaatg gcttgaagtt atgtgtttaa atctgctcat 1500
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 55 gagcggccccc cctggacgc ctccttcata caaactgcct ttgcctggc cctacgacag 180
 56 tcgagacgac ttccccctcc gaaaaacagc ctctgaaccc aacttgaag tgcgttcaag 240
 57 gctaaaacag aagggtggctg agcggagaag cagtccttc ctgcgtcgca aggatggac 300
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 61 gcaccgagcc ctccctctgg acagtccttc caaccatgc agcctctaca cgtctccctc 540
 62 tctgccaac atctccctag ggctgcaggg cacggtaact gtcaccaact cacacccac 600
 63 tgccctcccg aagctgtcga cacagcagga ggccgagagg cagggccctcc agtccctgcg 660
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 84 gctggagctg gccttcaagg tggctgcagg agagctcaag aatggatttg ccatcatccg 1920
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Input Set : N:\Crf3\RULE60\09502945.raw
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95	gcccttggat gaggcagtct tgcatggaaaa gcccaacatc aacgcagtgg ccacgctaga	2580
96	gaaagtcatc gagatccaga gcaaacactg gagctgtgtg cagaagttcg ccgctggct	2640
97	ggccgggtcc ctgcgagggg cccaaggcagg tgagaccgaa gaagccgaaa tgtgaacgcc	2700
98	atggccttgc tggtgggtgg ggcgaacag gcccaagctg cggcagcccg ggaacacagc	2760
99	cccaggccgg cagaggagcc catggagcag gaggctgccccc tgtgacgccccc cgccccccat	2820
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126	<223> OTHER INFORMATION:	
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130	<223> OTHER INFORMATION:	
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Input Set : N:\Crf3\RULE60\09502945.raw
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150	tgtatggagag agaaagcatc naaagcttct ggaagcaatc agttcccttg atggaaagaa	180
151	taggcggaaa ttggctgana ggtctgaggc tagtctgaag gtgtcagagt tcaatgtcag	240
152	ttctgaagga tcaggagaaa agctggctt tgcaaatctg cttgagccctg ttaaaaacttc	300
153	atcttctttg gccactgtga aaaagcaact gagtagagtc anatcaaaga anacagtgg	360
154	gttacctctg aacaaagaag agattgaacg gatccacaga gaatagcatt caataaaacg	420
155	cacaagtctt ctccaaatgg gaccctgtcg tcctgaagaa ccggcaggca gagcagctgg	480
156	ttttccccc ggagaaagag gagccagcca ttgctcccat tgaacatgtg ctcagtggt	540
157	ggaaggcaag aactccctg gaggagaaa ttttcaaccc cctccataag aacaaggcagc	600
158	cagtgcacaga ccctttactg accccctgtgg aaaaggccctc tctccgagcc atgagcctag	660
159	aaggaggaaa gatgcgacga gcagagcttc agagggctcg ggctctgcag tcctactatg	720
160	angccaaggc tcgaagagag aagaaaatcn aaagttaaaa gtatcacaaa gtcgtgaaga	780
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162	ccgcactaga agaacgaaaga aaagaggaaa gaaggaggag gagaaagaag aagaacaagg	900
163	agaagaagaa agaagaaggaa agaaggagaa gaaaagaagg agaagaggaa aaggaagaagg	960
164	gagaagaaaa aggagaagga aaaggaaaag aaggagaaga aagaagaact aagaagaagg	1020
165	agaggaagaa taagaaggaa agaagaaaga aaaaagtnaa agaagaagaa agaaggaaga	1080
166	aggaaaagaag aggaagaact nagaagaaga aagaggagga aagaagaaaag aagaataagg	1140
167	aacnagaaag aaggagaaga aagaataaga agaggaagaa gaaaaagaag aaaagaagaa	1200
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178	tccggcata aggtggattt tctgattgaa aatgtatgcag agaaggacta tctctatgtat	180
179	gtgctgcgaa tgttaccacca gaccatggac gtggccgtgc tcgtgggaga cctgaagctg	240
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182	cgtctggacc gtctgcaccc cgaaggcctc ggcctgagtg tgcgtgggtt cctggagtt	420
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186	ctgatccccg tggaaatctc tcctgtatgg ccccttactt ggcagttatgt ggatcgttt	660
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188	gagaagaagg tcttcatcg cctggtaggc tcccgaggcc ttggctgcag cattccagc	780
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191	ctggatcaca aggaggctgt aaatgtctg aaaaatagcc gcagcctgac catctccatt	960
192	gtagctgcag ctggccggga gctgttcatg acagaccggg agcggctggc agaggcgcgg	1020
193	cagcgtgagc tgcagcggca ggagcttctc atgcagaagc ggctggcgtat ggagtcac	1080
194	aagatccctcc aggaggcagca ggagatggag cggcaaagga gaaaagaaat tgcccagaag	1140

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198	gaacctgagc tcgagccccgc agatgacctg gatggaggca cggaggagca gggagagcag	1380
199	gatttccgga aatatgagga aggcttgac ccctactcta tggccatccc agaggcagatc	1440
200	atggggaaagg atgtcccgct cctacgcattc aagaaggagg gatcctttaga cctggccctg	1500
201	gaaggcgggtg tggactcccc cattggaaag gtggctgttt ctgctgtgt tgagcgggaa	1560
202	gctgctgagc ggcattgggg cattgtgaaa ggggacgaga tcatggcaat caacggcaag	1620
203	attgtgacag actacaccct ggctgaggct gacgctgccc tgcagaaggc ctggaatcag	1680
204	ggcggggact ggatcgaccc tgggggtgcc gtctggccccc caaaggagta tgacgatgag	1740
205	ctgaccccttct tgctgaagtc caaaaggggaa aaccaaattc acgcgtttagg aaacagttag	1800
206	ctccggccccc acctcgtaa cacaaggctt cggaccagcc ttgagagagg ccacatgaca	1860
207	cacaccagat ggcattccctt ggacctgaat ctatcaccctt ggaatctcaa actcccttt	1920
208	gccctgaacc agggccagat aaggaacagc tcggggccact ttttgaagg ccaatgtgga	1980
209	gaaaaggggag cagccagccg tttggggagaa gatctcaagg atccagactc tcattccctt	2040
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211	taagaccctt ctggagtctc tctctccca tcccttcctt ctgcctctg ctctaattgc	2160
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223	gtgctgcaaa tgtaccacca gaccatggac gtggccgtgc tcgtgggaga cctgaagctg	240
224	gtcatcaatg aacccagccg tctgcctctt tttgatgcca ttcggccgct gatcccactg	300
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234	gaggtggat tggagatagg ggaccatgg tgcgaagtca atggcgttca cttctctaa	900
235	ctggatcaca aggaggctgt aaatgtgtt aaaaatagcc gcagcctgac catctccatt	960
236	gtagctgcaat ctggccggaa gctgttcatc acagaccggg agcggctggc agaggcgg	1020
237	cagcgtggc tgcagcggca ggagcttctc atgcagaagc ggctggcgat ggagtcac	1080
238	aagatcccttcc aggaggcagca ggagatgggg cggcaaaaggaa gaaaagaaat tgcccaaa	1140
239	gcagcagagg aaaaatgagag ataccgaaag gagatgaaac agattttaga ggagggaaag	1200
240	aagttaaga agcaatggga agaagactgg ggctcaaagg aacagctact cttgcctaaa	1260
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242	tgaggaaggc tttgaccctt actctatgtt cacccttccatc cagatcatgg ggaaggatgt	1380
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244	ctccccattt gggaaagggtgg tcgtttctgc tggatgtatgtt cggggagctg ctgagcggca	1500

RAW SEQUENCE LISTING ERROR SUMMARY
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:3; N Pos. 55,141,199,342,352,722,750,1058,1101,1144

VERIFICATION SUMMARY

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Input Set : N:\Crf3\RULE60\09502945.raw
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L:119 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:3,Line#:0
L:123 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:3,Line#:0
L:127 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:3,Line#:0
L:131 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:3,Line#:0
L:135 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:3,Line#:0
L:139 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:3,Line#:0
L:143 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:3,Line#:0
L:147 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:3,Line#:0
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M:341 Repeated in SeqNo=3